## SEQUENCE LISTING

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Kristensen, Peter

Jestin, Jean-Luc

Winter, Gregory

<120> Selection System

<130> 8039/1090

<140> 09/710,444

<141> 2000-11-20

<150> GB 9810223.9

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<150> GB 9810228.8

<151> 1998-05-13

<150> PCT/GB99/01526

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<170> PatentIn version 3.1

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Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu Gly Arg Gly Ala His
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ggcaccetca gaacggtace ceaeceteag aggceggetg ggcegeeace eteagag
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ggtggcggcc cagccggcct ttctgagggg tcgactatag aaggacgagg gcccagcgaa
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ggaggtggg taccccttc tgagggtgg
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ccaccetcag aagggggtac cccacctcct tegetgggcc etegteette tatagtegac
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ccctcagaaa ggccggctgg gccgccacc
                                                                     89
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gcgatggttg ttgtcattgt cggc
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aaaagaaacg caaagacacc acgg
                                                                     24
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cctcctgagt acggtgatac acc
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<400> 6

<212> DNA

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<220>
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attttcggtc atagccccct tattag
                                                                     26
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<220>
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caaacgggcg gccgcagact acaaggatga cgacgacaag gaaactgttg aaagttgttt
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agcaa
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<210> 11
<211> 51
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<400> 9

<212> DNA

<213> Artificial sequence

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<223> Synthetic PCT primer used to change codon usage in recombinant cl
       ones
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cccctcagaa aggccggctg ggccgccgcc agcattgaca ggaggttcag g
                                                                     51
<210> 12
<211> 52
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<220>
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      ones.
<220>
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<222> (1)..(52)
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       ones
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gaaggaggtg gggtacccgg ttccgagggt ggttccggtt ccggtgattt tg
                                                                     52
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ccctcggaac cggtacccca gctgcttcgt gggccc
                                                                     36
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<210> 14

<211> 47

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<213> Bacillus amyloliquefaciens
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ctggcggcgg cccagccggc cctgcacagg ttatcaacac gtttgac
                                                                    47
<210> 15
<211> 43
<212> DNA
<213> Bacillus amyloliquefaciens
<400> 15
ctcggaaccg gtacctctga tttttgtaaa ggtctgataa gcg
                                                                    43
<210> 16
<211> 44
<212> DNA
<213> Gallus gallus
<400> 16
ggcggcccag ccggcctttc tctctctgac gaggacttca aggc
                                                                    44
<210> 17
<211> 41
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<212> DNA

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cctcggaacc ggtaccgaag agtcctttct ccttcttgag g
                                                                     41
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<220>
<221> misc_feature
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<223> Synthetic PCR primer used for library construction
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tacgccaagc ttgcatgc
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ctgcacctgg gccatgg
<210> 20
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<223> Synthetic PCR primer used for library construction.
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<400> 20
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gattacgcca agctttg

17

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<210> 21
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<211> 126

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<213> Erwinia chrysanthemi

<220>

<221> misc\_feature

<223> n at positions 23, 24, 29, 55, 56, 81, 97, 101, and 102 can be G,
A, T or C

<220>

<221> misc\_feature

<222> (23)..(23)

<223> n at position 23 can be G, A, T or C

<220>

<221> misc\_feature

<222> (24)..(24)

<223> n at position 24 can be G, A, T or C

<220>

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<221> misc_feature
<222> (29)..(29)
<223> n at position 29 can be G, A, T or C
<220>
<221> misc_feature
<222> (55)..(55)
<223> n at position 55 can be G, A, T or C
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<222> (56)..(56)
<223> n at position 56 can be G, A, T or C
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<222> (81)..(81)
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<220>
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<222> (101)..(101)
<223> n at position 101 can be G, A, T or C
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<221> misc_feature
<222> (102)..(102)
<223> n at position 102 can be G, A, T or C
<400> 21
gattacgcca agcttgcatg cannddctnt dtcaaggaga cagtcataat garrnnbcta
                                                                     60
ttgsyaayrs yasyasyagb nttgttatta ctcsyanycv nncygdccat ggcccaggtg
                                                                    120
cagctg
                                                                    126
<210> 22
<211> 117
<212> DNA
<213> Bacteriophage M13mp18
<220>
<221> misc_feature
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<222> (18)..(18)
<223> Nucleotide at position 18 can be G, A, T or C.
<220>
<221> misc_feature
<222> (19)..(19)
<223> Nucleotide at position 19 can be G, A, T or C.
<220>
<221> misc_feature
<222> (20)..(20)
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<220>
<221> misc_feature
<222> (21)..(21)
<223> Nucleotide at position 21 can be G, A, T or C.
<400> 22
gattacgcca agctttgnnn ncttttttww ggagattttc aacrtgaraa rattattatt
                                                                     60
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csyaattsyt ttagttsyts ytttctwtgy ggyccagccg gccatggccc aggtgca

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<210> 23
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<213> Artificial sequence

<220>

<223> Synthetic PCR primer used for vector construction.

<400> 23

ctttatgctt ccggctcg

18

<211> 17

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic PCR primer for library construction.

<220>

<221> misc\_feature

<222> (1)..(17)

<223> Synthetic PCT primer for library construction

<400> 24

cggccccatt cagatcc

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<210> 25
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<223> Randomized E. chrysanthemi pelB sequence
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aagcttgcat gcaaattcta tdtcaaggag acagttataa tgaaatacct
<210> 26
<211> 50
<212> DNA
<213> Artificial sequence
<220>
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<223> Randomized E. chrysanthemi pelB sequence.

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<220>
<221> misc_feature
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<223> Randomized E. chrysanthemi pelB sequence
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<221> misc_feature
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<223> n at position 14 can be G, A, T or C.
<220>
<221> misc_feature
<222> (15)..(15)
<223> n at position 15 can be G, A, T or C.
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<221> misc_feature
<222> (20)..(20)
<223> n at position 20 can be G, A, T or C.
<220>
<221> misc_feature
<222> (45)..(45)
<223> n at position 45 can be G, A, T or C.
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<220>
<221> misc_feature
<222> (46)..(46)
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<223> n at position 46 can be G, A, T or C.

<400> 26

aagcttgcat gcannddctn tdtcaaggag acagtcataa tgarrnnbct

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<210> 27

<211> 50

<212> DNA

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<223> Randomized E. chrysanthemi pelB sequence.

<220>

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<222> (1)..(50)

<223> Randomized E. chrysanthemi pelB sequence

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aagcttgcat gcagcatctc tdgcaaggag acagtcataa tgaagacgct

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<223> Randomized E. chrysanthemi pelB sequence
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aagcttgcat gcacgggctg tdtcaaggag acagtcataa tgagagggct
                                                                     50
<210> 29
<211> 50
<212> DNA
<213> Artificial sequence
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<223> Randomized E. chrysanthemi pelB sequence.

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<221> misc_feature
<222> (1)..(50)
<223> Randomized E. chrysanthemi pelB sequence
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aagcttgcat gcaccagctc tdtcaaggag acagtcataa tgaggcggct
<210> 30
<211> 55
<212> DNA
<213> Artificial sequence
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<223> Randomized E. chrysanthemi pelB sequence.
<220>
<221> misc_feature
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<400> 30

<222> (1)..(55)

attectaacg geageegetg gattgttatt actegeggee cageeggeea tggee

<223> Randomized E. chrysanthemi pelB sequence

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<210> 31
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<222> (38)..(38)
<223> n at position 38 can be G, A, T or C.
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<221> misc\_feature

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<222> (42)..(42)
<223> n at position 42 can be G, A, T or C.
<220>
<221> misc_feature
<222> (43)..(43)
<223> n at position 43 can be G, A, T or C.
<400> 31
attgsyaayr syasyasyag bnttgttatt actcsyanyc vnncygdcca tggcc
<210> 32
<211> 55
<212> DNA
<213> Artificial sequence
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<223> Randomized E. chrysanthemi pelB sequence.
<220>
<221> misc_feature
<222> (1)..(55)
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<223> Randomized E. chrysanthemi pelB sequence

```
<400> 32
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attgcyaatg gtactgtyag gattgttatt actcccaccc ggtccgtcca tggcc

55

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<210> 33
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<211> 55

<212> DNA

<213> Artificial sequence

<220>

<223> Randomized E. chrysanthemi pelB sequence.

<220>

<221> misc\_feature

<222> (1)..(55)

<223> Randomized E. chrysanthemi pelB sequence

<400> 33

attgcyaatg ctagtgcyag ggttgttatt actcccaatc gcgccggcca tggcc

55

<210> 34

<211> 54

<212> DNA

<213> Artificial sequence

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<221> misc_feature
<222> (43)..(43)
<223> n at position 43 can be G, A, T or C.
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<221> misc_feature
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<400> 34

<210> 35 <211> 22 <212> PRT

<213> Erwinia chrysanthemi

<400> 35

Met Lys Tyr Leu Leu Pro Thr Ala Ala Gly Leu Leu Leu Ala

1 5 10 15

Ala Gln Pro Ala Met Ala

20

<210> 36

<211> 22

<212> PRT

<213> Artificial sequence

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<221> MISC\_FEATURE

<222> (1)..(22)

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Met Arg Gly Leu Ala Met Leu Val Ala Gly Gly Pro Ile Ala Pro Ala 1 5 10 15

Gln Pro Ala Met Ala

20

<210> 38

<211> 23

<212> PRT

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<220>

<221> MISC\_FEATURE

<222> (1)..(23)

<223> Randomized E. chrysanthemi pelB sequence

<400> 38

Met Arg Arg Leu Val Pro Ile Thr Ala Ala Val Gly Leu Leu Ala Pro

1 5 10 15

```
Pro Thr Gln Pro Ala Met Ala
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20

<210> 39

<211> 50

<212> DNA

<213> Artificial sequence

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<223> Randomized bacteriophage M13 g3 sequence.

<220>

<221> misc\_feature

<222> (1)..(50)

<223> Randomized bacteriophage M13 g3 sequence

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aagctttgga cgcttttttt tggagatttt caacgtgaaa aaattattat

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<210> 40

<211> 50

<212> DNA

<213> Artificial sequence

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<223> Randomized bacteriophage M13 g3 sequence.
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<222> (9)..(9)
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<220>
<221> misc_feature
<222> (1)..(50)
<223> Randomized bacteriophage M13 g3 sequence
<220>
<221> misc_feature
<222> (10)..(10)
<223> n at position 10 is can be G, A, t or C.
<220>
<221> misc_feature
<222> (11)..(11)
<223> n at position 11 is can be G, A, t or C.
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<220>

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aagctttgnn nncttttttw wggagatttt caacrtgara arattattat
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<223> Randomized bacteriophage M13 g3 sequence.
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aagctttggg gcctttttt aggagatttt caacatgaga agattattat
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<210> 42

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<211> 50
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<221> misc_feature
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<210> 43
<211> 50
<212> DNA
<213> Artificial sequence
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<223> Randomized bacteriophage M13 g3 sequence.
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<221> misc_feature
<222> (1)..(50)
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<400> 43
tcsyaattsy tttagttsyt sytttctwtg yggyccagcc ggccatggcc
                                                                     50
<210> 44
<211> 50
<212> DNA
<213> Artificial sequence
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<223> Randomized bacteriophage M13 g3 sequence.
<220>
<221> misc_feature
<222> (1)..(50)
<223> Randomized bacteriophage M13 g3 sequence
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tcctaattcc tttagttgtt gctttctatg tggtccagcc ggccatggcc
                                                                    50
<210> 45
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<223> Randomized bacteriophage M13 g3 sequence

<211> 22

<212> PRT

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<213> Artificial sequence
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<223> Randomized bacteriophage M13 g3 sequence.
<220>
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<222> (1)..(22)
<223> Randomized bacteriophage M13 g3 sequence
<400> 45
Met Lys Lys Leu Leu Phe Ala Ile Pro Leu Val Val Pro Phe Tyr Ala
1
                5
                                   10
                                                       15
Ala Gln Pro Ala Met Ala
            20
<210> 46
<211> 22
<212> PRT
<213> Artificial sequence
<220>
<223> Randomized bacteriophage M13 g3 sequence.
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<220>
<221> MISC_FEATURE
<222> (1)..(22)
<223> Randomized bacteriophage M13 g3 sequence
<400> 46
Met Arg Arg Leu Leu Leu Ala Pro Pro Val Ala Val Pro Phe Tyr Val
                5
                                    10
                                                        15
Val Gln Pro Ala Met Ala
            20
<210> 47
<211>
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<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic oligonucleotide primer used as a substrate for Stoffel
       fragment of Thermus aquaticus DNA polymerase I.
<220>
<221> misc_feature
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<222> (1)..(18)
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<223> Synthetic oligonucleotide primer used as substrate for Stoffel fr agment of Thermus aquaticus DNA polymerase I

<400> 47

tttcgcaaga tgtggcgt

18

- <210> 48
- <211> 12
- <212> DNA
- <213> Artificial sequence

<220>

<223> Synthetic oligonucleotide primer used as a substrate for Thermus aquaticus DNA polymerase I.

<220>

- <221> misc\_feature
- <222> (1)..(12)

<223> Synthetic primer used as substrate for Stoffel fragment of Thermu s aquaticus DNA polymerase I

<400> 48

gcgaagatgt gg

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<211> 30
<212> DNA
<213> Artificial sequence
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<223> Synthetic oligonucleotide primer used as a substrate for Thermus
       aquaticus DNA polymerase I.
<220>
<221> misc_feature
<222> (1)..(30)
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       uaticus DNA polymerase I
<400> 49
aaatacaaca ataaaacgcc acatcttgcg
                                                                     30
<210> 50
<211> 20
<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic oligonucleotide sequence insert containing PstI restric
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<210> 49

tion site and frame shift for H102A mutant barnase fusion construct fused to p3 gene of phage fd-3.

<220>

<221> misc\_feature

<222> (1)..(20)

<223> Synthetic oligonucleotide sequence insert containing PstI restric tion site and frame shift for H102A mutant barnase fusion construct fused to p3 gene of phage fd-3.

<400> 50

ctgcaggcgg tgcggccgca

20

<210> 51

<211> 24

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotide used for random priming.

<220>

<221> misc\_feature

<222> (1)..(24)

<223> Synthetic oligonucleotide used for random priming

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<220>
<221> misc_feature
<222> (19)..(19)
<223> n at position 19 can be G, A, T or C.
<220>
<221> misc_feature
<222> (20)..(20)
<223> n at position 20 can be G, A, T or C.
<220>
<221> misc_feature
<222> (21)..(21)
<223> n at position 21 can be G, A, T or C.
<220>
<221> misc_feature
<222> (22)..(22)
<223> n at position 22 can be G, A, T or C.
<220>
<221> misc_feature
<222> (23)..(23)
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<223> n at position 23 can be G, A, T or C.
<220>
<221> misc_feature
<222> (24)..(24)
<223> n at position 24 can be G, A, T or C.
<400> 51
gagcctgcag agctcaggnn nnnn
                                                                     24
<210> 52
<211> 23
<212> DNA
<213> Artificial sequence
<220>
<223> Synthetic PCR primer used to re-amplify randomply amplified E. co
       li genomic DNA sequence.
<220>
<221> misc_feature
<222> (1)..(23)
<223> Synthetic PCR primer used to re-amplify randomly amplified E. col
       i genomic DNA sequences.
```

```
<400> 52
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cgtgcgagcc tgcagagctc agg

23

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<210> 53
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<211> 45

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(45)

<223> Barstar binding barnase-p3 fusion insert

<400> 53

Leu Gln Ser Ser Gly Asp Cys Val Ile Ser Asp Thr Cys Ile Ala Gly

1 5 10 15

Met Ala Glu Ala Ala Cys Glu Glu Lys Phe Ser Ser Gln Asn Val

Gly Leu Thr Ile Thr Val Thr Pro Cys Leu Ser Ser Ala

35 40 45

<210> 54

<211> 44

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 54

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

```
Pro Ser Ser Ala Thr Ile His Cys Leu Ser Ser Ala
```

35

<210> 55

<211> 40

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(40)

<223> Barstar binding barnase-p3 fusion insert

<400> 55

Leu Gln Ser Ser Gly Asp Ser Ala Gly Cys Lys Asn Met Thr Gly Gly

1 5 10 15

Arg Leu Tyr Ala His Thr Leu Glu Ala Ile Ile Pro Gly Phe Ala Val

```
Ser Ala Pro Ala Cys Glu Pro Ala
```

40

<210> 56

<211> 33

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 56

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His

1 5

10

15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro

```
Ala
```

<210> 57

<211> 44

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 57

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

```
Pro Ser Ser Ala Thr Val Gln Cys Leu Ser Ser Ala
```

<210> 58

<211> 41

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(41)

<223> Barstar binding barnase-p3 fusion insert

<400> 58

Leu Gln Ser Ser Gly Lys Ile Val Gln Ala Gly Ala Asn Ile Gln Asp

1 5 10 15

Gly Cys Ile Met His Gly Tyr Cys Asp Thr Asp Thr Ile Val Gly Glu

```
Asn Gly His Ile Gly Leu Ser Ser Ala
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<210> 59

<211> 45

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(45)

<223> Barstar binding barnase-p3 fusion insert

<400> 59

Leu Gln Ser Ser Gly Val Cys Val Ile Ser Asp Thr Cys Ile Ala Gly

1 5 10 15

Thr Ala Glu Ala Ala Cys Glu Glu Lys Phe Ser Ser Gln Asn Val

```
Gly His Thr Ile Thr Glu Thr Pro Cys Leu Ser Ser Ala
```

45

35

<220>

<220>

<210>	60
<211>	44
<212>	PRT
<213>	Artificial sequence

<223> Barstar binding barnase-p3 fusion insert.

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 60

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu
20 25 30

```
Pro Ser Ser Ala Thr Ile Gln Cys Leu Ser Ser Ala
```

35

<210> 61

<211> 53

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(53)

<223> Barstar binding barnase-p3 fusion insert

<400> 61

Leu Gln Ser Ser Gly Gln Asp Ser Gln Arg Glu His Ala Ser His Thr

1 5

10

15

Ala Glu Asp Asp Cys Glu Asp Gln Thr Arg Ile His Gln His Ile Arg

```
Glu Val Asp Phe Val Asp Thr Pro Gln Glu Val Asp Asp Cys Arg Ala
```

35 40 45

Ala Leu Ser Ser Ala

50

<210> 62

<211> 33

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 62

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His

1 5 10 15

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro

20

25

30

Ala

<210> 63

<211> 9

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(9)

<223> Barstar binding barnase-p3 fusion insert

<400> 63

Leu Gln Ser Ser Gly Val Arg Pro Ala

1

```
<210> 64
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<211> 44

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(44)

<223> Barstar binding barnase-p3 fusion insert

<400> 64

Leu Gln Ser Ser Gly Cys Gly Ser Ser Gly Ser Ser Ile Asn Cys Leu

1 5 10 15

Pro Cys Gly Ala Thr Ser Arg Gly Thr Ser Pro Leu Ala Ser Gly Leu

20 25 30

Pro Ser Ser Ala Thr Ile Gln Cys Leu Ser Ser Ala

```
<210> 65
<211> 30
<212> PRT
<213> Artificial sequence

<220>
<223> Barstar binding barnase-p3 fusion insert.

<220>
<221> MISC_FEATURE
<222> (1)..(30)
<223> Barstar binding barnase-p3 fusion insert
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Leu Gln Ser Ser Gly Thr Glu Val Asp Arg Gly Asn Gln Gln His Asp

1 5 10 15

Thr Asn Asp Arg Asp Phe Thr His Thr Pro Leu Ser Ser Ala
20 25 30

<210> 66 <211> 36

<400> 65

<212> PRT

```
<213> Artificial sequence
<220>
<223> Barstar binding barnase-p3 fusion insert.
<220>
<221> MISC_FEATURE
<222> (1)..(36)
<223> Barstar binding barnase-p3 fusion insert
<400> 66
Leu Gln Ser Ser Gly Val Ala Gln Gly Ser Ser Ala Ser Val Asp Val
1
                5
                                    10
                                                        15
Thr Ala Thr Asn Ala Val Leu Ser Ala Asp Ser Leu Ser Leu Gly Gly
            20
                                25
                                                    30
Gly Glu Pro Ala
        35
<210> 67
<211> 19
```

<212> PRT

```
<213> Artificial sequence
<220>
<223> Barstar binding barnase-p3 fusion insert.
<220>
<221> MISC_FEATURE
<222> (1)..(19)
<223> Barstar binding barnase-p3 fusion insert
<400> 67
Leu Gln Ser Ser Gly Gly Ala Val Ala Val Thr Pro Gly Pro Val Leu
1
                5
                                    10
                                                       15
Ser Ser Ala
<210> 68
<211> 18
<212> PRT
<213> Artificial sequence
<220>
<223> Barstar binding barnase-p3 fusion insert.
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<220> <221> MISC\_FEATURE <222> (1)..(18) <223> Barstar binding barnase-p3 fusion insert <400> 68 Leu Gln Ser Ser Gly His Cys Arg Gly Lys Pro Val Leu Cys Thr His 5 10 15 Thr Ala <210> 69 <211> 9 <212> PRT <213> Artificial sequence <220> <223> Barstar binding barnase-p3 fusion insert. <220> <221> MISC\_FEATURE

<222> (1)..(9)

```
<223> Barstar binding barnase-p3 fusion insert
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<400> 69

Leu Gln Ser Ser Gly Val Arg Pro Ala

1 5

<210> 70

<211> 36

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 70

Leu Gln Ser Ser Gly Glu Pro Ala Pro Ala His Glu Ala Lys Pro Thr

1

Glu Ala Pro Val Ala Lys Ala Glu Ala Lys Pro Glu Thr Pro Ala His

20

25

30

Leu Ser Ser Ala

35

<210> 71

<211> 33

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(33)

<223> Barstar binding barnase-p3 fusion insert

<400> 71

Leu Gln Ser Ser Gly Cys Val Arg Leu Lys Arg Thr Ser Val Asn His

1

5

10

Gln Pro Asp Ala Trp Pro Glu Pro His Leu Lys Ala Ala Cys Glu Pro

20

25

30

Ala

<210> 72

<211> 36

<212> PRT

<213> Artificial sequence

<220>

<223> Barstar binding barnase-p3 fusion insert.

<220>

<221> MISC\_FEATURE

<222> (1)..(36)

<223> Barstar binding barnase-p3 fusion insert

<400> 72

Leu Gln Ser Ser Gly Val Val Asp Trp Ala Lys Met Arg Glu Ile Ala

1

Asp Ser Ile Gly Ala Tyr Leu Phe Val Asp Met Ala His Val Ala Ala
20 25 30

Leu Ser Ser Ala

35

<210> 73

<211> 117

<212> DNA

<213> Artificial sequence

<220>

<223> Vector pK1 polylinker sequence.

<220>

<221> misc\_feature

<222> (1)..(117)

<223> Vector pK1 polylinker sequence

<400> 73

aatgctggcg gcggcccagc cggcctttct gaggggtcga ctatagaagg acgaggggcc

60

cacgaaggag gtgggtacc cggttccgag ggtggttccg gttccggtga ttttgat

<210> 74

<211> 39

<212> PRT

<213> Artificial sequence

<220>

<223> Polypeptide encoded by pK1 vector polylinker sequence.

<220>

<221> MISC\_FEATURE

<222> (1)..(39)

<223> Polypeptide encoded by pK1 vector polylinker sequence

<400> 74

Asn Ala Gly Gly Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu

1 5 10 15

Gly Arg Gly Ala His Glu Gly Gly Gly Val Pro Gly Ser Glu Gly Gly
20 25 30

Ser Gly Ser Gly Asp Phe Asp

```
<210> 75
<211> 117
<212> DNA
<213> Artificial sequence
<220>
<223> Vector pK2 polylinker sequence.
<220>
<221> misc_feature
<222> (1)..(117)
<223> vector pK2 polylinker sequence
<400> 75
aatgctggcg gcggcccagc cggcctttct gaggggtcga ctatagaagg acgagggccc
                                                                     60
acgaagcagc tggggtaccg gttccgaggg tggttccggt tccggtgatt ttgatta
                                                                    117
<210> 76
<211> 39
<212> PRT
<213> Artificial sequence
<220>
<223> Polypeptide sequence encoded by vector pK2 polylinker region.
```

```
<220>
<221> MISC_FEATURE
<222> (1)..(39)
<223> Polypeptide sequence encoded by vector pK2 polylinker region.
<220>
<221> MISC_FEATURE
<222> (38)..(38)
<223> X represents a TGA stop codon
<220>
<221> MISC_FEATURE
<222> (36)..(36)
<223> X represents a stop codon (TGA)
<400> 76
Asn Ala Gly Gly Gly Pro Ala Gly Leu Ser Glu Gly Ser Thr Ile Glu
                5
                                    10
                                                        15
```

Gly Arg Gly Pro Thr Lys Gln Leu Gly Tyr Arg Phe Arg Gly Trp Phe

```
Arg Phe Arg Xaa Phe Xaa Leu
```

<210> 77

<211> 35

<212> DNA

<213> Artificial sequence

<220>

<223> Sequence of the junction region between Barnase and p3 in recombinant fusion vector fd-3.

<220>

<221> misc\_feature

<222> (1)..(35)

<223> Sequence of the junction region between Barnase and p3 in recombinant fusion vector fd-3.

<400> 77

atcagactgc aggcggtgcg gccgcagaaa ctgtt

35

<210> 78

<211> 11

<212> PRT

<213> Artificial sequence <220> <223> Amino acid sequence about the junction of Barnase and p3 coding r egions of recombinant fusion vector fd-3. <400> 78 Ile Arg Leu Gln Ala Ala Ala Ala Glu Thr Val 5 10 <210> 79 <211> 4 <212> PRT <213> 'Artificial sequence <220> <223> Factor Xa protease cleavage sequence. <220> <221> MISC\_FEATURE <222> (1)..(1) <223> X can be either Ile or Leu.

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<221> MISC_FEATURE

<222> (1)..(4)

<223> Factor Xa proteolytic cleavage site.

<400> 79

Xaa Glu Gly Arg

??

??

(continued...)
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(continued...)